



STQ / GAU 1644
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PATENT
Our Docket: P-IX 3536

NOV 09 2000

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

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In re application of:)
Huse and Wu)
Serial No.: 09/339,922)
Filed: June 24, 1999)
For: ANTI- $\alpha_v\beta_3$ RECOMBINANT)
HUMAN ANTIBODIES,)
NUCLEIC ACIDS ENCODING)
SAME AND METHODS OF USE)

Group Art Unit: 1644

Examiner: P. Gambel

I hereby certify that this correspondence is being deposited with
the United States Postal Service as first class mail in an envelope
addressed to: Box Sequence, Commissioner for Patents,
Washington, D.C. 20231, on October 31, 2000.

By Deborah L. Cadena
Deborah L. Cadena, Reg. No. 44,048

October 31, 2000
Date of Signature

TECH CENTER 1600/2900

NOV 13 2000

BOX SEQUENCE

Commissioner for Patents
Washington, D.C. 20231

Sir:

TRANSMITTAL

Applicants respectfully request the Examiner's
consideration of the following:

- X 1) a Communication regarding sequences;
- X 2) a computer readable form of the sequence listing;
- X 3) a paper copy of the sequence listing, pages 1
through 35;
- X 4) a statement under 37 CFR §1.825(a) and (b); and
- X 5) a return receipt post card.


c

Inventors: Huse and Wu
Serial No.: 09/339,922
Filed: June 24, 1999
Page 2

No fee is deemed necessary in connection with the filing of this document. However, if any fee is required, authorization is hereby given to charge the amount of any such fee to Deposit Account No. 03-0370.

Respectfully submitted,

Date: October 31, 2000



Deborah L. Cadena
Registration No. 44,048
Telephone: (858) 535-9001
Facsimile: (858) 535-8949

CAMPBELL & FLORES LLP
4370 La Jolla Village Drive
7th Floor
San Diego, California 92122
USPTO CUSTOMER NO. 23601

C



has dlc PIX 3536
**UNITED STATES DEPARTMENT OF COMMERCE
Patent and Trademark Office**

Address: COMMISSIONER OF PATENTS AND TRADEMARKS
Washington, D.C. 20231
Nov 09 2000

APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCK
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013801
CAMPBELL & FLORES LLP
4370 LA JOLLA VILLAGE DRIVE
7TH FLOOR
SAN DIEGO CA 92161

PM11/1003

CAMPBELL EXAMINER

ART UNIT

PAPER NUMBER

DOCKETED

*Sequences Due
11-3-00*

DATE MAILED:

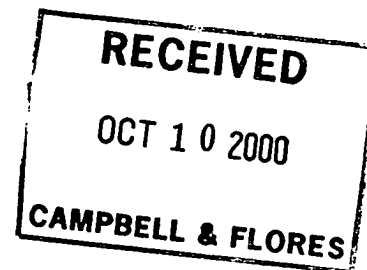
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Please find below and/or attached an Office communication concerning this application or proceeding.

Commissioner of Patents and Trademarks



COPY

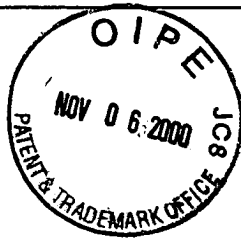


UNITED STATES DEPARTMENT OF COMMERCE
Patent and Trademark Office
COMMISSIONER OF PATENTS AND TRADEMARKS
Washington, D.C. 20231

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TECH CENTER 1600/2900

091339922

SERIAL NUMBER	FILING DATE	FIRST NAMED APPLICANT	ATTORNEY	SECRET NO.



APPLICANT'S
COPY

EXAMINER	
Mary Tung	
ART UNIT	PAPER NUMBER
1644	

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DATE MAILED:

NOV 13 2000

Please find below a communication from the EXAMINER in charge of this application
Commissioner of Patents and Trademarks
TECH CENTER 1600/2900

This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 CAR 1.821(a)(1) and (a)(2). However, the CRF submission filed 10/12/99 fails to comply with the requirements of 37 CAR 1.821 through 1.825 for the reason(s) set forth on the attached Notice To Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures. Applicant must comply with the requirements of the sequence rules (37 CAR 1.821 - 1.825) before the application can be examined under 35 U.S.C. §§ 131 and 132.

The disk submitted contains errors, as set forth in the attached "Raw Sequence Listing Error Report." Applicants are required to submit a computer readable disk and a substitute paper copy of the sequences according to the attached "Notice to Comply with the Sequence Rules." Applicant is reminded of the sequence rules which require a submission for all sequences of more than 9 nucleotides or 3 amino acids (see 37 CAR 1.821-1.825) and is also requested to carefully review the submitted specification for any and all sequences which require compliance with the rules.

Any inquiry concerning this communication should be directed to Examiner Phillip Gambel, Art Unit 1644, whose telephone number is (703) 308-3997.

Any inquiry of a general nature or relating to the status of this application should be directed to the Group receptionist whose telephone number is (703) 308-0196.

Applicant is given ONE MONTH, or THIRTY DAYS, whichever is longer, from the mailing date of this letter within which to comply with the sequence rules, 37 CAR 1.821 - 1.825. Failure to comply with these requirements will result in ABANDONMENT of the application under 37 CAR 1.821(g). Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 CAR 1.136(a). In no case may an applicant extend the period for reply beyond the SIX MONTH statutory period. Direct the reply to the undersigned. Applicant is requested to return a copy of the attached Notice to Comply with the reply.

PHILLIP GAMBEL
Phillip Gambel, Ph.D.
Primary Examiner
Art Unit 1644
Technology Center 1600
October 2, 2000

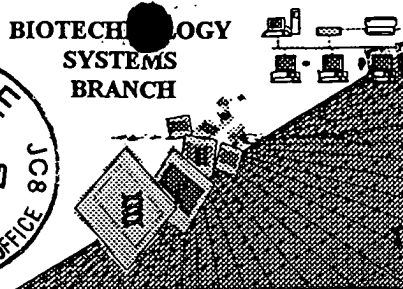
2

Handled

RAW SEQUENCE LISTING
ERROR REPORT



BIOTECHNOLOGY
SYSTEMS
BRANCH



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The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

NOV 13 2000

TECH CENTER 1600/2900

Application Serial Number: 09/339,922

Source: 1644

Date Processed by STIC: 9/26/2000

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR FURTHER INFORMATION, PLEASE TELEPHONE MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

C



P. Gambel

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NOV 09 2000

1644

TECH CENTER 1600/2900

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/339,922

DATE: 09/26/2000
 TIME: 08:04:08

Input Set : A:\Ix35361.app
 Output Set: N:\CRF3\09262000\I339922.raw

Does Not Comply
 Corrected Diskette Needed

3 <110> APPLICANT: Huse, William D.
 4 Wu, Herren
 6 <120> TITLE OF INVENTION: Anti-AlphaV Beta3 Recombinant Human Antibodies, Nucleic
 7 Acids Encoding Same and Methods of Use
 9 <130> FILE REFERENCE: P-IX 3536
 11 <140> CURRENT APPLICATION NUMBER: US 09/339,922
 12 <141> CURRENT FILING DATE: 1999-06-24
 14 <160> NUMBER OF SEQ ID NOS: 112
 16 <170> SOFTWARE: PatentIn Ver. 2.1
 18 <210> SEQ ID NO: 1
 19 <211> LENGTH: 351
 20 <212> TYPE: DNA
 21 <213> ORGANISM: Artificial Sequence
 23 <220> FEATURE:
 24 <221> NAME/KEY: CDS
 25 <222> LOCATION: (1)..(351)
 27 <220> FEATURE:
 28 <223> OTHER INFORMATION: Description of Artificial Sequence: grafted
 29 antibody variable region
 31 <400> SEQUENCE: 1
 32 cag gtg cag ctg gtg gag tct ggg gga ggc gtt gtg cag cct gga agg 48
 33 Gln Val Gln Leu Val Glu Ser Gly Gly Val Val Gln Pro Gly Arg
 34 1 5 10 15
 36 tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc agt agc tat 96
 37 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
 38 20 25 30
 40 gac atg tct tgg gtt cgc cag gct ccg ggc aag ggt ctg gag tgg gtc 144
 41 Asp Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 42 35 40 45
 44 gca aaa gtt agt agt ggt ggt ggt agc acc tac tat tta gac act gtg 192
 45 Ala Lys Val Ser Ser Gly Gly Gly Ser Thr Tyr Tyr Leu Asp Thr Val
 46 50 55 60
 48 cag ggc cga ttc acc atc tcc aga gac aat agt aag aac acc cta tac 240
 49 Gln Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 50 65 70 75 80
 52 ctg caa atg aac tct ctg aga gcc gag gac aca gcc gtg tat tac tgt 288
 53 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 54 85 90 95
 56 gca aga cat aac tac ggc agt ttt gct tac tgg ggc caa ggg act aca 336
 57 Ala Arg His Asn Tyr Gly Ser Phe Ala Tyr Trp Gly Gln Gly Thr Thr
 58 100 105 110
 60 gtg act gtt tct agt 351
 61 Val Thr Val Ser Ser
 62 115
 65 <210> SEQ ID NO: 2
 66 <211> LENGTH: 117
 67 <212> TYPE: PRT

see
 Apr 2, 3, 5
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NOV 13 2000

TECH CENTER 1600/2900

RAW SEQUENCE LISTING

DATE: 09/26/2000

PATENT APPLICATION: US/09/339,922

TIME: 08:04:08

Input Set : A:\Ix35361.app

Output Set: N:\CRF3\09262000\I339922.raw

68 <213> ORGANISM: Artificial Sequence
 69 <220> FEATURE:
 69 <223> OTHER INFORMATION: Description of Artificial Sequence: grafted
 72 <400> SEQUENCE: 2
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 74 1 5 10 15
 76 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Tyr
 77 20 25 30
 79 Asp Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
 80 35 40 45
 82 Ala Lys Val Ser Ser Gly Gly Gly Ser Thr Tyr Tyr Leu Asp Thr Val
 83 50 55 60
 85 Gln Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
 86 65 70 75 80
 88 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
 89 85 90 95
 91 Ala Arg His Asn Tyr Gly Ser Phe Ala Tyr Trp Gly Gln Gly Thr Thr
 92 100 105 110
 94 Val Thr Val Ser Ser
 95 115
 99 <210> SEQ ID NO: 3
 100 <211> LENGTH: 321
 101 <212> TYPE: DNA
 102 <213> ORGANISM: Artificial Sequence
 104 <220> FEATURE:
 105 <221> NAME/KEY: CDS
 106 <222> LOCATION: (1)..(321)
 108 <220> FEATURE:
 109 <223> OTHER INFORMATION: Description of Artificial Sequence: grafted
 110 antibody variable region
 112 <400> SEQUENCE: 3
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 114 Glu Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Leu Ser Pro Gly
 115 1 5 10 15
 117 gaa agg gcg act ctt tcc tgc cag gcc agc caa agt att agc aac cac 96
 118 Glu Arg Ala Thr Leu Ser Cys Gln Ala Ser Gln Ser Ile Ser Asn His
 119 20 25 30
 121 cta cac tgg tat caa caa agg cct ggt caa gcc cca agg ctt ctc atc 144
 122 Leu His Trp Tyr Gln Gln Arg Pro Gly Gln Ala Pro Arg Leu Leu Ile
 123 35 40 45
 125 aag tat cgt tcc cag tcc atc tct ggg atc ccc gcc agg ttc agt ggc 192
 126 Lys Tyr Arg Ser Gln Ser Ile Ser Gly Ile Pro Ala Arg Phe Ser Gly
 127 50 55 60
 129 agt gga tca ggg aca gat ttc acc ctc act atc tcc agt ctg gag cct 240
 130 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro
 131 65 70 75 80
 133 gaa gat ttt gca gtc tat tac tgt caa cag agt ggc agc tgg cct cac 288
 134 Glu Asp Phe Ala Val Tyr Tyr Cys Gln Ser Gly Ser Trp Pro His
 135 85 90 95

per 1.823 of new sequence rules,
 <220> is mandatory
 whenever <221>,
 <222>, or <223>
 is present.

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/339,922

DATE: 09/26/2000
TIME: 08:04:08

Input Set : A:\Ix35361.app
Output Set : N:\CRF3\09262000\I339922.raw

321

137 acg ttc gga ggg ggg acc aag gtg gaa att aag
138 Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
139 100
142 <210> SEQ ID NO: 4
143 <211> LENGTH: 107
144 <212> TYPE: PRT
145 <213> ORGANISM: Artificial Sequence

W---> 146 <220> FEATURE:
146 <223> OTHER INFORMATION: Description of Artificial Sequence: grafted

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151 1 5 10 15
153 Glu Arg Ala Thr Leu Ser Cys Gln Ala Ser Gln Ser Ile Ser Asn His
154 20 25 30
156 Leu His Trp Tyr Gln Gln Arg Pro Gly Gln Ala Pro Arg Leu Leu Ile
157 35 40 45
159 Lys Tyr Arg Ser Gln Ser Ile Ser Gly Ile Pro Ala Arg Phe Ser Gly
160 50 55 60
162 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Glu Pro
163 65 70 75 80
165 Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Ser Gly Ser Trp Pro His
166 85 90 95
168 Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
169 100 105

173 <210> SEQ ID NO: 5
174 <211> LENGTH: 351
175 <212> TYPE: DNA
176 <213> ORGANISM: Mus musculus
178 <220> FEATURE:
179 <221> NAME/KEY: CDS
180 <222> LOCATION: (1)..(351)

182 <400> SEQUENCE: 5
183 gaa gtg cag ctg gtg gag tct ggg gga ggc tta gtg aag cct gga agg 48
184 Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Lys Pro Gly Arg 15
185 1 5 10 15
187 tcc ctg aga ctc tcc tgt gca gcc tct gga ttc gct ttc agt agc tat 96
188 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Ala Phe Ser Ser Tyr 30
189 20 25 30
191 gac atg tct tgg gtt cgc cag att ccg gag aag agg ctg gag tgg gtc 144
192 Asp Met Ser Trp Val Arg Gln Ile Pro Glu Lys Arg Leu Glu Trp Val 45
193 35 40 45
195 gca aaa gtt agt agt ggt ggt agc acc tac tat tta gac act gtg 192
196 Ala Lys Val Ser Ser Gly Gly Gly Ser Thr Tyr Leu Asp Thr Val 60
197 50 55 60
199 cag ggc cga ttc acc atc tcc aga gac aat gcc aag aac acc cta tac 240
200 Gln Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Tyr 70
201 65 70 75 80
203 ctg caa atg agc agt ctg aac tct gag gac aca gcc atg tat tac tgt 288
204 Leu Gln Met Ser Ser Leu Asn Ser Glu Asp Thr Ala Met Tyr Tyr Cys 75

insert
2237
at
beginning
of each
line



RAW SEQUENCE LISTING
PATENT APPLICATION:

US/09/339,922

DATE: 09/26/2000
TIME: 08:04:08

Input Set : A:\I35361.app
Output Set: N:\CRF3\09262000\I339922.raw

205
207 gca aga cat aac tac ggc agt ttt gct tac tgg ggc caa ggg act ctg 336
208 Ala Arg His Asn Tyr Gly Ser Phe Ala Tyr Trp Gly Gln Gly Thr Leu 110 351
209 100
211 gtc act gtc tct gca
212 Val Thr Val Ser Ala
213 115
216 <210> SEQ ID NO: 6
217 <211> LENGTH: 117
218 <212> TYPE: PRT
219 <213> ORGANISM: Mus musculus
221 <400> SEQUENCE: 6
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223 1 5 10 15
225 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Ala Phe Ser Ser Tyr 30
226 20 25 30
228 Asp Met Ser Trp Val Arg Gln Ile Pro Glu Lys Arg Leu Glu Trp Val 45
229 35 40 45
231 Ala Lys Val Ser Ser Gly Gly Gly Ser Thr Tyr Tyr Leu Asp Thr Val 60
232 50 55 60
234 Gln Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Tyr 75 80
235 65 70 75
237 Leu Gln Met Ser Ser Leu Asn Ser Glu Asp Thr Ala Met Tyr Tyr Cys 95
238 85 90
240 Ala Arg His Asn Tyr Gly Ser Phe Ala Tyr Trp Gly Gln Gly Thr Leu 110
241 100 105
243 Val Thr Val Ser Ala
244 115
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249 <211> LENGTH: 321
250 <212> TYPE: DNA
251 <213> ORGANISM: Mus musculus
253 <220> FEATURE:
254 <221> NAME/KEY: CDS
255 <222> LOCATION: (1)..(321)
257 <400> SEQUENCE: 7
258 gat att gtg cta act cag tct cca gcc acc ctg tct gtg aca cca gga 48
259 Asp Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Val Thr Pro Gly 15
260 1 5 10
262 gat agc gtc agt ctt tcc tgc cag gcc agc caa agt att agc aac cac 96
263 Asp Ser Val Ser Leu Ser Cys Gln Ala Ser Gln Ser Ile Ser Asn His 30
264 20 25 30
266 cta cac tgg tat caa caa aaa tca cat gag tct cca agg ctt ctc atc 144
267 Leu His Trp Tyr Gln Gln Lys Ser His Glu Ser Pro Arg Leu Leu Ile 45
268 35 40
270 aag tat cgt tcc cag tcc atc tct ggg atc ccc tcc agg ttc agt ggc 192
271 Lys Tyr Arg Ser Gln Ser Ile Ser Gly Ile Pro Ser Arg Phe Ser Gly 60
272 50 55
274 agt gga tca ggg aca gat ttc gct ctc agt atc aac agt gtg gag act 240



2

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/339,922

DATE: 09/26/2000

TIME: 08:04:08

Input Set : A:\Ix35361.app

Output Set: N:\CRF3\09262000\I339922.raw

275 Ser Gly Ser Gly Thr Asp Phe Ala Leu Ser Ile Asn Ser Val Glu Thr
 276 65 70 75 80
 278 gaa gat ttt gga atg tat ttc tgt caa cag agt ggc agc tgg cct cac 288
 279 Glu Asp Phe Gly Met Tyr Phe Cys Gln Gln Ser Gly Ser Trp Pro His
 280 85 90 95
 282 acg ttc gga ggg ggg acc aag ctg gaa att aag 321
 283 Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
 284 100 105
 287 <210> SEQ ID NO: 8
 288 <211> LENGTH: 107
 289 <212> TYPE: PRT
 290 <213> ORGANISM: Mus musculus
 292 <400> SEQUENCE: 8
 293 Asp Ile Val Leu Thr Gln Ser Pro Ala Thr Leu Ser Val Thr Pro Gly
 294 1 5 10 15
 296 Asp Ser Val Ser Leu Ser Cys Gln Ala Ser Gln Ser Ile Ser Asn His
 297 20 25 30
 299 Leu His Trp Tyr Gln Gln Lys Ser His Glu Ser Pro Arg Leu Leu Ile
 300 35 40 45
 302 Lys Tyr Arg Ser Gln Ser Ile Ser Gly Ile Pro Ser Arg Phe Ser Gly
 303 50 55 60
 305 Ser Gly Ser Gly Thr Asp Phe Ala Leu Ser Ile Asn Ser Val Glu Thr
 306 65 70 75 80
 308 Glu Asp Phe Gly Met Tyr Phe Cys Gln Gln Ser Gly Ser Trp Pro His
 309 85 90 95
 311 Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
 312 100 105
 316 <210> SEQ ID NO: 9
 317 <211> LENGTH: 84
 318 <212> TYPE: DNA
 319 <213> ORGANISM: Artificial Sequence
 321 <220> FEATURE:
 322 <223> OTHER INFORMATION: Description of Artificial Sequence:
 323 oligonucleotide
 325 <400> SEQUENCE: 9
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 327 tcctgtgcag cctctggatt cacc 84
 330 <210> SEQ ID NO: 10
 331 <211> LENGTH: 84
 332 <212> TYPE: DNA
 333 <213> ORGANISM: Artificial Sequence
 335 <220> FEATURE:
 336 <223> OTHER INFORMATION: Description of Artificial Sequence:
 337 oligonucleotide
 339 <400> SEQUENCE: 10
 340 aacttttgcg acccactcca gacccttgcc cggagcctgg cgaaccaag acatgtcata 60
 341 gctactgaag gtgaatccag aggc 84
 344 <210> SEQ ID NO: 11
 345 <211> LENGTH: 87

FYI:

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/339,922

DATE: 09/26/2000

TIME: 08:04:09

Input Set : A:\Ix35361.app

Output Set: N:\CRF3\09262000\I339922.raw

L:69 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:146 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:618 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:638 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:668 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:886 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:918 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:950 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:982 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1014 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:1078 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:1238 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:1791 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:1858 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1890 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1922 M:258 W: Mandatory Feature missing, <220> FEATURE:

